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detection showed a linear response to log increases in the target level

over a 105-fold range, permitting the detn. of target level within an

order of magnitude. The assay showed .apprx. 109-fold discrimination over

Chlamydia pneumoniae (TWAR) rRNA. High levels of cultured

albicans, Escherichia coli, Staphylococcus aureus, or Neisseria gonorrhoeae had no detectable effect on assay background or the ability to

detect a single elementary body.

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OREF 126:13605a,13608a

TI Detection of EBV early RNA (EBER-1) in parotid pleomorphic adenomas: a

novel observation utilizing ***ligation*** -dependent PCR AU Brandwein, Margaret; Li, Hongbo; Zhang, David Y.

CS Lillian and Henry M. Stratton-Hans Popper Department of Pathology,

University of New York, NY, USA

SO International Congress Series (1996), 1114(Head and Neck Cancer: Advances

in Basic Research), 401-409

CODEN: EXMDA4; ISSN: 0531-5131

PB Elsevier

DT Journal

LA English

AB Very little is known regarding the initiation and promotion of salivary

neoplasia. We utilized the recently introduced ***ligation***

-dependent polymerase chain reaction (LD-PCR) to detect viral RNAs. This

technique employs two ***capture*** *** probes*** for the isolation

of target RNA. A third probe contains a complementary region to the $\,$

target sequence at each end, and a generic linker region for PCR primer

binding. This probe becomes circularized upon

hybridization to

the target and forms a covalently linked circular probe by incubation with

T4 DNA ****ligase*** . The circularized probe sequence serves as a

template for Taq polymerase. A novelty of this assay is to amplify by PCR

the probe sequence rather than the target sequence. This allows for the

facile identification of RNA by PCR without the need for the

transcriptase step. We studied nine cases of snap-frozen tissue from

parotid gland and six pleomorphic adenomas by LD-PCR for the presence of $% \left(1\right) =\left(1\right) \left(1\right)$

EBV early RNA (EBER-1). EBER-1 was identified in six of eight parotid

tissue samples and four of six pleomorphic adenomas. Although the $\ensuremath{\mathsf{PCR}}$

technique does not allow for localization within tissue (viral sequences

within tumor cells vs. circulating lymphocytes), the identification of

 $\ensuremath{\mathsf{EBER}}\xspace\text{-}1$ in these cases does indicate that active transcription of a

latency-assocd. viral RNA is common in the parotid gland. This may have

implications on salivary tumorigenesis.

OSC. G: 3 THERE ARE 3 CAPLUS RECORDS THAT CITE THIS RECORD (3 CITINGS)

L14 ANSWER 3 OF 6 CAPLUS COPYRIGHT 2010 ACS on STN AN 1995:230412 CAPLUS < LOGINI D:: 20100420> >

DN 122:179542

OREF 122:32745a,32748a

TI A rapid, reliable method for detection of known point mutations:

point-EXACCT

AU Somers, Veerle A. M. C.; Moerkerk, Peter T. M.; Murtagh, James J., Jr.;

Thunnissen, Frederik B. J. M.

CS Department Pathology, University Limburg, Maastricht, Neth.

SO Nucleic Acids Research (1994), 22(22), 4840-1 CODEN: NARHAD; ISSN: 0305-1048

PB Oxford University Press

DT Journal

LA English

AB Point mutations in the human genome play a central role in tumorigenesis.

Several methods are available for detection of known point mutations. The $\,$

detection format is based on an extension of the $\ensuremath{\mathsf{EXACCT}}$ procedure. In

 $\mbox{short: after exonuclease digestion, polymerase chain reaction} \label{eq:contraction} fragments$

are detd. by ***hybridization*** with a capture and a detection probe

complementary to sequences near the 3' end of the antisense fragment. The

capture *** probe*** bears a biotin residue and the other probe

digoxigenin. After *** hybridization*** the PCR product hybrids are

captured in streptavidin-coated microtiter plates and detected with

labeled anti-digoxigenin antibody. For the detection of known point

mutations this procedure was extended by using after the capture step the

*** ligation*** of a mutation-specific *** capture***
*** probe* **

with adjacent detection probe (Point-EXACCT). Point-EXACCT requires

considerably less time and effort than other techniques used

detection of known point mutations. This method is easily automated

permitting rapid screening of tissue banks with multiple probes to

individual base substitutions, deletions or addns. The simplicity of

Point-EXACCT makes it a highly promising method for the detection of known

AN 1994:70885 CAPLUS < LOGINID::20100420>>

OSC.G 15 THERE ARE 15 CAPLUS RECORDS THAT CITE THIS DN 120:70885 OREF 120:12639a,12642a RECORD (15 CITINGS) TI Chlamydiae probes for use in solution phase sandwich L14 ANSWER 4 OF 6 CAPLUS COPYRIGHT 2010 ACS on STN * * * hybridization* * * AN 1994:70887 CAPLUS << LOGINI D::20100420>> assavs DN 120:70887 IN Sanchez-Pescador, Ray: Besemer, Diana J.: Urdea, Michael OREF 120:12639a,12642a S. TI Cytomegalovirus (CMV) probes for use in solution phase PA Chiron Corp., USA sandwich SO PCT Int. Appl., 84 pp. * * * hybridization* * * assays CODEN: PLXXD2 IN Kolberg, Janice A.; Shen, Lu Ping; Urdea, Michael S. DT Patent PA Chiron Corp., USA LA English SO PCT Int. Appl., 71 pp. FAN. CNT 1 CODEN: PIXXD2 KIND DATE PATENT NO. APPLICATION NO DT Patent LA English --------- ------FAN. CNT 1 Pl WO 9313221 A1 19930708 WO 1992-US11035 PATENT NO. KIND DATE APPLICATION NO. 19921222 W: AU, CA, JP, KR DATE ---- ------RW: AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LU, MC, Pl WO 9313227 A1 19930708 WO 1992-US11170 NL. PT. SE 19921222 AU 9334672 A 19930728 AU 1993-34672 W: CA, JP, KR 19921222 RW: AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LU, MC, EP 726963 A1 19960821 EP 1993-903387 NL. PT. SE 19921222 EP 625214 A1 19941123 EP 1993-902723 R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LI, LU, MC, 19921222 NL, PT, SE R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LI, LU, MC, US 5618674 A 19970408 US 1995-479487 NL. PT. SE 19950607 US 5407795 A 19950418 US 1993-138608 PRAI US 1991-813587 A 19911223 A 19921222 WO 1992-US11035 19931015 ASSIGNMENT HISTORY FOR US PATENT AVAILABLE IN LSUS PRAI US 1991-813590 A 19911223 WO 1992-US11170 W 19921222 DISPLAY FORMAT ASSIGNMENT HISTORY FOR US PATENT AVAILABLE IN LSUS AB The title probes, i.e. amplifier probe and ***capture*** DISPLAY FORMAT * * * probe* * * AB The title probes, i.e. amplifier or ***capture*** , comprise a first segment with nucleotide sequence " * * probes* * * . substantially comprises a nucleotide sequence complementary to a complementary to a segment of Chlamydiae plasmid DNA and segment of CMV nucleic a second segment acid and a nucleotide sequence complementary to a segment with nucleotide sequence substantially complementary to an of nucleotide oliaonucleotide sequence of an amplifier multimer or a capture solid phase, multimer or an oligonucleotide bound to a solid phase, resp. Thus, a resp. Thus, a comb-type polynucleotide having 15 branch sites and side comb-type polynucleotide having 15 branch sites and side chain extensions chain extensions having 3 labeled probe binding sites was prepd. as an having 3 labeled probe binding sites was synthesized and amplifier multimer. used as a labeled OMV amplifier and ***capture*** *** probes*** multimer. The amplifier and ***capture*** probes*** are (contg., in addn. to *** hybridized*** with sample, the formed complexes are sequences complementary to CMV sequences, a 5' extension captured by complementary to the amplifier multimer or a downstream sequence of oligonucleotide-bound solid phase, and the captured CTTCTTTGGAGAAAGTGGTG complexes are *** hybridized* ** with the oligonucleotide multimer and complementary to an immobilized oligonucleotide, resp.) were complementary used along with the amplifier multimer and capture solid phase in a labeled oligonucleotide for Chlamydiae detection. OSC.G 7 THERE ARE 7 CAPLUS RECORDS THAT CITE THIS * * * hybridization* * * assay of CMV. RECORD (7 CITINGS) OSC.G 10 THERE ARE 10 CAPLUS RECORDS THAT CITE THIS RECONT 1 THERE ARE 1 CITED REFERENCES AVAILABLE FOR RECORD (10 CITINGS) THIS RECORD RE.ONT 2 THERE ARE 2 CITED REFERENCES AVAILABLE FOR ALL CITATIONS AVAILABLE IN THE RE FORMAT THIS RECORD L14 ANSWER 6 OF 6 CAPLUS COPYRIGHT 2010 ACS on STN ALL CITATIONS AVAILABLE IN THE RE FORMAT AN 1993:663642 CAPLUS << LOGINID::20100420>> L14 ANSWER 5 OF 6 CAPLUS COPYRIGHT 2010 ACS on STN DN 119:263642

point mutations.

OREF 119:46973a,46976a TI A transcriptionally amplified DNA probe assay with ligatable probes and immunochemical detection AU Carpenter, William R.; Schutzbank, Ted E.; Tevere, Vincent J.; Tocyloski, Kenneth R.: Dattagupta, Nanibushan: Yeung, Kwok K. CS Diagn. Div., Miles Inc., Tarrytown, NY, 10591, USA SO Clinical Chemistry (Washington, DC, United States) (1993), 39(9), 1934-8 CODEN: CLCHAU; ISSN: 0009-9147 DT Journal LA English AB Transcriptionally amplified DNA probes are valuable tools in the development of sensitive nucleic acid-based diagnostic assays. Here the authors describe a model assay using a novel oligonucleotide hairpin probe that encodes a T7 RNA polymerase promoter. The hairpin probe and an adjacently *** hybridizing *** biotinylated *** capture *** *** probe*** were *** hybridized*** to target DNA and the duplex was captured onto streptavidin-coated magnetic particles. After *** ligation*** of the immobilized probes, which served to specificity, the hairpin probe was transcribed by T7 RNA polymerase. The amplified RNA product was *** hybridized*** to the *capture*** *** probe*** and bound to the streptavidin-coated magnetic particles. The immobilized heteroduplex was detected with an antibodyalk. phosphatase conjugate specific for DNA: RNA hybrids, and the chemiluminescent substrate adamantyl-1,2-dioxetane Ph phosphate. Ten attomoles of target DNA could be detected in a background of 5 .mu.g of unrelated DNA. The chemiluminescent immunoassay was as sensitive as radioactive detection of specific product after gel electrophoresis. OSC.G 5 THERE ARE 5 CAPLUS RECORDS THAT CITE THIS RECORD (5 CITINGS) => log y COST IN U.S. DOLLARS SINCE FILE ENTRY SESSION FULL ESTIMATED COST 97.83 98.05 DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS) SINCE

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